

	Total	N-Clones (208F-FE-8)	T-Clones (FE8-208F)
Number of sequenced cDNA clones	1257	669	588
Number of individual sequences	823	416	407
Sequence analysis			
Known genes (nr/Genbank)	427	207	220
Expressed Sequence Tags (dbest)	303	161	142
No similarity in data bases (new)	93	48	45
Expression analysis: Reverse Northern Analysis/conventional Northern Blot			
Differentially expressed	393	225	168
Known genes	244	126	118
Expressed sequence tags	104	74	30
New sequences	45	25	20
Not differentially expressed	194	86	108
Not detectable in expression analysis	236	105	131

FIG. 1

FIG. 2

Genes that are adjusted down by H-Ras+transformation				Genes that are adjusted up by H-Ras+transformation			
Sequence Identity (Gebhardt/Bailey)	Species identity	Accession number	Extent of change	Sequence Identity (Gebhardt/Bailey)	Species identity	Accession number	Extent of change
3'-5'-cyclic AMP phosphodiesterase	r Z2867	1	>100 N1	AKAP-KC (A kinase anchor protein)	r	AH03276	1 16.1
AP-1 repressor	m A015140	1	38.0 R	B51 (lect receptor ligand)	m	DB0515	1 5.2
CAM-dependent protein kinase type II	r M2492	1	>100 R	C-htrax-1	h	W00574	1 17.0
CSP-1 (colon stimulating factor-1)	r M8161	2	5.6 N2,R	Cainodulin-dependent protein kinase II-delta	m	X67677	1 12.5
Gas-6	m X9846	1	24.0 R	Cyclodenylase I	r	J05072	1 8.1
Guanine nucleotide-binding protein G-s alpha	r M2673	1	3.6 N3	Cyclophilin-B-binding protein 1	r	U03288	1 90.7
T- ³² P-ATP-interacting protein	m V058864	1	38.6 N4	Cyclophilin-B-specific immunophilin	r	U28830	1 8.3
IKK-complex-associated protein (IKAP)	h A001195	1	8.6 R	FKBP1 (T-cell specific immunophilin)	m	U56599	1 68.2
M2CKS	m M60714	2	3.3 N5	FLIP (FLICE-like inhibitor protein)	m	U97016	2 >100
MST2 kinase	r A0001539	2	21.6 R	GRF-H4-binding protein R455	h	U72206	1 32.1
Myo-inositol monophosphatase (IMP)	r U04338	1	44.5 N6	GRF-binding protein R455	r	AC0702935	1 >100
P5 protein	ha X26278	1	3.4 R	JAK1 protein-tyrosine kinase 1	r	A000556	1 55.0
Protein-kinase-like protein (PKLP)	r 115354	2	>100 N7,R	MAP-kinase phosphatase (CG21)	r	AC013144	1 27.9
Phosphatidylinositol 3-kinase PI(0) beta	h S57334	1	>100 N8	MAP-kinase phosphorylated 67 kDa protein	r	MB0367	1 98.2
Phosphatidylinositol 3-kinase PI(0)	m U55772	1	65.5 N9,R	Phosphatase 2A h56	r	LA2373	2 50.6
Protein tyrosine phosphatase delta (PTPdelta)	m D13803	1	1.9 R	RKA kinase	r	V15748	1 19.9
ROK alpha	r J38881	1	26.1 N10	RPS2	r	J14453	1 >100
Serum inducible kinase (SKN)	m M81653	1	>100 N11,R	Rap1B GRF-binding protein	r	U07195	1 21.0
SH3 binding protein (SAB)	h A0003047	1	3.5 R	Ras-GTPase-activating protein	m	A0001927	1 9.9
				RhoC	m	X00338	2 6.7
				SB1 phosphatase	h	U33181	1 27.1
				Spctry 2 (SPY2)	h	AC03943	2 11.60
				TMG31	m	U44088	1 2.7
				Syntapse phosphatase Ta-2&	r	D59222	1 12.2

Nuclear Proteins (Transcription Factors, DNA Processing Enzymes)									
ANRNP nucleoprotein	h	M03002	2	>100	N12				
ATP-dependent RNA helicase	m	U6590	1	8.9	N13	Alpha-1-proteinase		1	2.4
BRO1 (ribribo homolog)	m	S8108	1	13.1	N14	BRCA1-associated RING domain protein (Bard1)	m	3.5	123
CCA/T/enhancer binding (C/EBP Gamma)	r	X04403	1	16.6	N15	cdk-1-like kinase (Clk)	m	13.1	124
Cdc21	m	D05089	1	3.9	R	CDN-1 (cap endonuclease-1)	m	11.1	125
Centromeric protein CENPC	m	U03113	1	39.2	N16,R	Fra-1 (forselated antigen 1)	m	11.1	125
Chromosome-associated polypeptide C (CAP-C)	h	A01987	1	9.6	R	Histone acetyltransferase (G95)	h	>100	T66,R
DNA repair protein Rad50	m	S05699	1	5.1	R	Hsp65 nucleolar protein	h	2.7	127
DNAbinding protein Rb50	m	U66887	1	3.4	N17,R	IAB/C (lamina associated polypeptide 1C)	m	1.6	128
DSX transcription factor	h	U7163	1	9.6	N18	Myo-binding Protein (176)	m	1.6	129
ETC TEA domain containing transcription factor	m	D05063	1	7.4	N19	NP-1 transcriptional coactivator	m	0.9	130
Gp130 binding protein	h	U7824	1	41.7	N20	p100 transcriptional coactivator	h	0.8	131
HE4 retinoblastoma-associated protein	h	A017790	1	3.9	N21,R	PBP2b2	m	0.4	132
Histone H1	h	A015812	2	>100	N22,R	Rb (retinoblastoma protein)	m	0.3	133
K1-67 antigen	m	Z89560	2	5.8	R	SA-1 (nuclear antigen)	m	0.3	134,R
Lrp2 (Lambda associated polypeptide 2)	m	X02186	1	>100	N23,R				
Mouse zinc finger protein	f	U78113	4	2100	R				
MTF33 (X-linked transcriptional activator)	m	D05210	1	5.6	N25				
Nuclear astacinian G2NA	h	S16673	1	3.6	R				
Nucleoporin 155	h	U7989	1	31.9	R				
Poly(A)-ribose) glycohydrolase (hARG)	m	A007558	1	15.2	N26				
Rad4 transcription factor	m	A019557	1	2.4	R				
Single strand DNA-binding protein	m	U85141	2	64.9	R				
STR51 transcription factor	h	A007048	1	4.9	R				
Topoisomerase I	f	U64715	1	1.8	N27				
Topoisomerase II	m	D00661	1	20.1	R				
	r	Z19352	3	2.1	R				
Protein Processing, Protein Transport and Protein-folding Molecule									
26S proteasome subunit p55	h	A003103	1	3.5	N28	Anionopeptidase PAP2	m	5.6	R
GAP43/endothelin	m	S8916	1	2.2	R	Chaperonin containing TCP-1 epsilon (CT)	m	2.2	T55,R
heat shock protein 105	m	D7016	1	15.1	N29	Exportin-a	h	48.5	336
Heat shock protein 90	a	X15183	1	4.8	N30,R	GP215	m	2.1	337

FIG. 2A

NC-I-160 (Golgi apparatus substrate protein)		beta5		Translocation protein-1	
I	008136	1	2..3	R	RATP (theres sas, ubiquitin-specific protease)
I	035355	1	56.0	N31	Importin alpha Q1
I	081127	1	>100	R32	NP92 (mitochondrial processing peptidase beta)
					Ran-GTPase
					Sec61
					Sortilin (sortilin)
					Translation initiation factor 3
Metabolic Enzymes, Transporters and Ion Channels					
r	S61167	4	12.7	R	472B (integral membrane glycoprotein)
r	X52625	2	12.7	R	ABC transporter M0AT1
r	J04637	1	37.8	K33	Aryl-CoA oxidase 1
m	X61112	1	6.3	R	Aldehyde dehydrogenase
m	W97436	1	1.8	N34	Angiogenesis synthase
m	555199	1	56.7	R	Anti-clotting-lyase
m	176713	1	>100	N35	Baclofen hydrolase
r	X73328	2	4.7	R36	ClC-6 (chloride channel)
r	M88751	1	18.8	N37	Fattyacyl pyrophosphate synthetase
r	D78014	2	2.3	R	Glutathione 6-phosphate dehydrogenase
r	M61632	3	10.4	R	Glutathione reductase
r	X1848	1	2.5	R	GIVr-1 (leukemia virus receptor 1)
r	X1220	1	5.3	R	GN1 (monooxybutyrate transporter)
m	G24024	1	12.3	N38	Mitochondrial trifunctional protein
r	U90516	1	6.2	N39	Monoubiquitin-encase (NEV)
r	W65374	2	31.8	N40	NP21 protein
					Phospholipase mutant type B
					Stenoy-CoA desaturase 2
					Tanscript-PAF (pathway of arachidonate differentiation)
					Transcript-PAF (G1)
					WIF1 (Wingless-type MMTV integration site family member 1)

FIG. 2B

FIG. 2C

Cytoskeleton Components-Molecule Involved in Adhesion and Cell-Cell Interaction

ABP-280 (actin-binding protein/filamin)	h X53416	1	5.8	R	Arg3 (actin-related protein 3)	h AF06083	3	3.3	T49,R
Alpha-actinin	r YG6001	5	4.2	R	Calcium-binding protein 1pc2/LSP1/WIF34	m M89596	2	29.7	T00,R
Calmodulin	m X71557	1	11.7	R	Caponin	r D0575	1	5.2	R
Cadermin	r U84119	3	37.7	M1	CB44 9-glycoprotein	r M6185	1	17.0	T01,R
Cytohesin-1	r U07728	1	51.0	M2	Larabin receptor	m J05970	5	4.1	R
Gas-1	m X51528	1	10.4	R	Leukocyte adhesion protein p150,95	h T0053	2	5.2	R
HSP core fibrogranin (syndecan-2)	r M61687	1	61.9	N43,R	MAGE-B gene cluster	h D0153	2	15.3	T52
ImMAP microtubule associated protein	h X004434	1	26.9	N44	Mycobacterium regulatory light chain	r D1488	1	6.9	R
MC-2	r S79700	2	2.6	N45,R	TAL oncocytic gene	r U0095	2	1.9	T53
Pedopain	m YG6340	1	60.1	N46	Taymanin Beta 4	r X34033	1	2.4	T54,R
Podopain	r U96549	1	9.4	R					
Rydorcan	r S81668	6	27.7	N47,R					
Tropocortin 4	r T00169	1	7.8	N48,R					
Tropo-2/clusterin	r M4723	1	39.4	N49					
Vimentin	r X22352	1	1.6	R					

Extracellular Proteins

Collagen alpha1	r Z78779	34	22.3	R	MMP-1 (Collagenase)	r M62516	19	>100	T55,R
Ory61 (immediate-early gene)	m M2490	4	10.0	N50,R	Myr3 (Siremyyan 1)	r X0561	2	32.3	T56,R
Enacrin/Nidogen	m X4194	14	35.8	N51	Myr10 (Siremyyan 2)	m X0563	12	33.3	R
Fibrillin-1 [Fn1]	m U2293	1	3.3	R	Mab-1	r X17035	2	2.4	T57,R
Fibronectin	r X5506	25	110.0	N52	Tastin	m X7890	1	8.9	T58
FIP-2	m M00642	2	49.4	N53					
Follicular-related protein: TSC-36	r U66664	5	2.0	N54,R					
Laminin B1	m M5525	1	5.0	R					
Lysyl oxidase	r U11338	14	9.2	R					
Lysyl oxidase-related protein (NS9-14)	h U99442	1	59.2	N55,R					
Mekaryocyte potentiating factor	m D6570	3	6.0	N56					
MGF (mat. cell growth factor)	m U44725	1	13.4	N57					
MGP-2 (Relatinase A)	r U55556	3	50.6	N58,R					
The combogedin 1	m M2270	25	42.5	R					
TIMP-2 (inhibitor of metalloproteinase 2)	r S23594	1	18.3	N59,R					



FIG. 2D

		Others	
MAG1 (anti-apoptotic gene)			
Xba-6 (activity and retrotranslitter-ind. gene 6)	h U38357	3.1	NG6 Annexin IV
Antifolin	r AK03091	1	B-cell receptor associated protein 37 (BAP 37)
ATP-dependent metalloendopeptidase Psh1	m S4728	2	B2-prion protein p32
C8B20 (GAP-binding protein)	m A900330	21.3	BESC-1 (breast cancer suppressor candidate 1)
Collapsin-2	h X44157	2	B1 (similar to Lysyl hydroxylase 1cctcm 3)
DCC-2196 Phosphoprotein	c U8240	10.2	C29 keratin-1 Related
E124 (p3 responsive gene)	r U55177	7.4	Calmodulin (RCM)
eIF-4E1 protein synthesis initiation factor	m U1151	5.5	EB1 19-kDa-2-binding protein homolog (Nip3)
H411 precursor	m X6653	3.9	Flt3L53 activated in colon tumors
Intron-1 induced gene	ha A006670	>100	Glycyl-tRNA synthetase
KIAA0415 (myeloblast)	h Y6121	1	HRH2216 rat fetal brain gene
KIAA0418 (myeloblast)	h D8876	16.3	Insulinoma Gene (rig)
KIAA0428 (myeloblast)	h D8018	33.8	RK359 (myeloblast)
KIAA0425 (myeloblast)	h D8708	4.8	KIAA0435 (myeloblast)
KIAA0429 (myeloblast)	h D8748	3.6	KIAA0510 (brain)
KIAA0432 (brain)	h A002230	1	KIAA0511 (brain)
L1 retronposon (LGP2)	r X43381	20.8	KIAA0525 (brain)
Lem3 (LINE 1 repetitive sequence)	f M6024	26.2	KIAA0544 (brain)
Yamagata gene	m A005433	1	KIAA0595 (brain)
Osteoglychin	m D31951	5	KIAA0597 (brain)
F5B382 (p3-binding protein)	m U9881	10.3	L11 protein (L1)
PERBP1	m D4556	1	L11 protein (L1)
PMEMC (maternal embryonic message gene 2)	m X95550	23.4	MAP domain protein
SF3B splicing factor	h L1187	2	Ma-1-pain large subunit (cbl1)
KDM2C	r X71764	10.4	Neutrin
Zinc-finger domain-containing protein	m U90554	1	OP9150 kDa oxygen regulated protein
ZNFX216 zinc finger protein	m A002071	6.7	PHD finger protein 2 (PHF2)
			Ries3 (rat spinocerebellar atrophy type 3 gene)
			Ser13 tRNA synthetase
			Syntaxin (tammexin VII)
			TIC22
			TGOL1 (tumor susceptibility protein)
			Tyrosine phosphatase-like protein 1a-2a; PTpS5
			U01652
			U72941 1 57.8 T39,R
			m X76633 2 42.8 T00,R
			h AF043384 1 2.8 T11,R
			h AF002612 1 6.9 T62
			m X18844 1 2.6 T63
			h AB113607 1 6.6 T64
			m AB11323 2 2.8 T64
			z AB11324 1 6.6 T65
			m AF041034 1 63.0 T65
			h AB02704 1 2.3 T66
			h D05510 1 12.0 R
			h AB10315 1 2.9 T66
			h U09513 1 1.6 T67
			h M13193 1 16.0 T68
			h AF061093 1 3.2 R
			h DS7717 1 6.0 R
			h AB002398 1 10.7 R
			h AB001891 1 2.5 R
			h AB011097 1 2.9 R
			h AB011116 1 9.4 R
			h AB011167 1 2.9 R
			h AB11169 1 4.2 R
			h AF058899 1 7.3 T59
			h U6116 1 >100 T00,R
			z AL03336 1 28.7 R
			z RM03956 1 1.8 R
			z U41853 1 9.3 T11
			h NM_005391 1 2.1 R
			h U12339 1 55.5 T72
			z U8136 3 2.4 R
			h L11319 1 2.2 R
			h AF09573 1 2.3 R
			m U5395 2 2.2 T73
			h U74393 11 74.9 T00,R



Expressed Sequence Tags (EST)		
Down-adjusted ESTs	Up-adjusted ESTs	
ESTAA003402	ESTAA76763 ESTAA262510 ESTAA03320 ESTAA067238 ESTAA086555 ESTAA22120 ESTAA22120 ESTAA15720 ESTAA15720 ESTAA16144 ESTAA16144 ESTAA170629 ESTAA170629 ESTAA180452 ESTAA180716 ESTAA1849586 ESTAA16936 ESTAA167114 ESTAA166316 ESTAA207146	ESTAA87476 ESTAA81418 ESTAA276806 ESTAA286338 ESTAA101936 ESTAA212329 ESTAA22531 ESTAA646031 ESTAA3270 ESTAA3270 ESTAA19551 ESTAA152120 ESTAA151531 ESTAA161533 ESTAA18552 ESTAA190173 ESTAA20370 ESTAA21212 ESTAA174560 ESTAA0719 ESTAA85918 ESTAA07513 ESTAA01125 ESTAA64710 ESTAA117802 ESTAA61247 ESTAA51788 ESTAA67143
25 New Sequences	New Sequences	
	20 New Sequences	

FIG. 2E

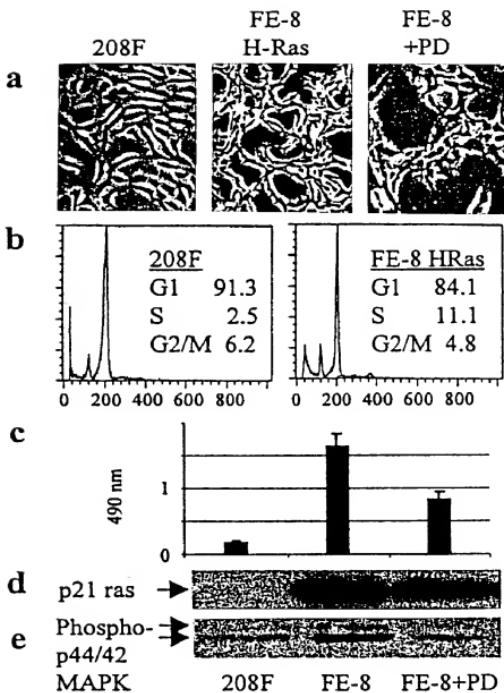
FIG. 3

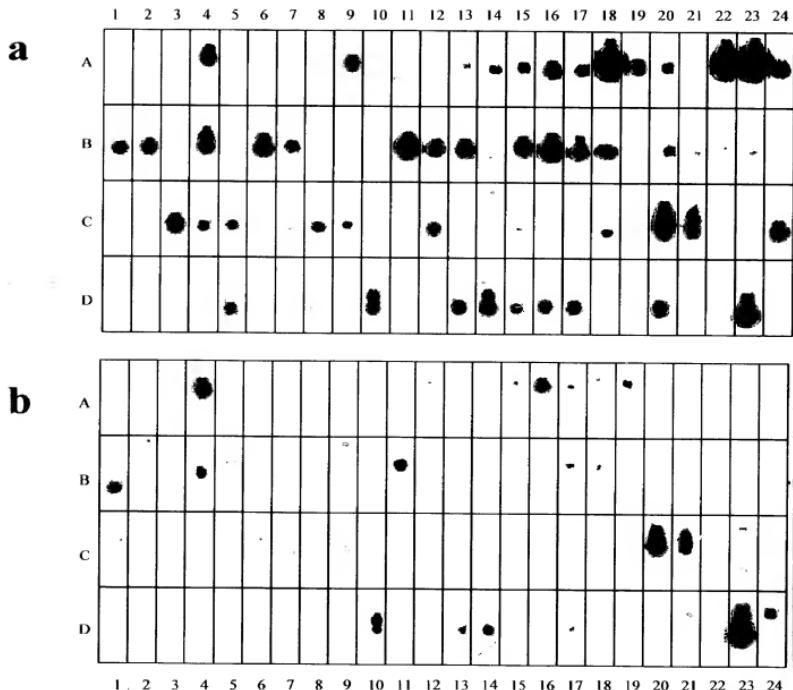
Sequence Identity (Genbank/ZBGU)	Expression Strength				Expression Strength			
	208F	F68	F68	+D	208F	F68	F68	+D
3-hydroxy-3-methylglutaryl coA synthase	+++	+	+++		+++	++	+++	++
ABP-280 (actin binding protein/filamin)	+++	++	+++		0	++	+	+
Alpha-actin	+++	+	+++		0	+++	++	++
Anticodon enzyme AOE372	+++	+	++		+	+++	++	++
APO26 (acetaminophen-binding protein)	++	0	++		0	+++	+	+
Cdc21	++	0	++		0	+++	+	+
Centromeric Protein CENPC (a)	+++	0	+++		0	++	0	+
Collagen alpha 1	+++	0	+++		0	+++	++	++
CSF-1 colony stimulating factor 1)	++	0	++		0	+++	0	+
DCC-5; p96 phosphoprotein	++	0	++		0	+++	+	+
FRS1 transcription factor	+++	+	++		0	+++	+	+
ETF transcription factor	+++	0	++		0	+++	+	+
Flaminacin	+++	+	++		0	+++	0	+
Follistatin-related protein; TSC36	++	+	+++		+	+++	+	+
GRP96/uncoplasmin	+++	+	++		0	++	0	+
Gα binding protein	+++	0	+		+	+++	++	++
Heat shock protein 90	++	0	++		0	+++	+	+
HSFC core fibrolycan (syndecan-2)	++	0	++		+	+++	+	+
Insectiron induced gene	+++	0	++		0	+++	+	+
IL receptor (CRF2)	+++	0	++		0	+++	+	+
Laminin B1	+++	0	++		0	+++	0	+
Lysyl oxidase	++	0	+		0	+++	+	+
Lysyl oxidase-related protein (WS9-14)	++	0	+		0	++	+	+
Mata gene	++	0	+		0	+++	+	+
Mon-2 (Galatinase A)	+	0	++		+	++	++	++
MRE13 (transcriptional activator)	++	+	++		0	+++	+	+
Nuclear autoantigen GS28A	++	0	++		0	+++	+	+
Osteoglycin	++	0	++		0	+++	+	+
P5 protein	+++	+	++		0	+++	+	+
P-adherin	++	0	++		0	+++	+	+
Protein-like protein (PPL2)	+++	0	++		0	++	+	+
Serin/arginine kinase (SAK)	+++	0	++		0	++	+	+
Stat1 transcription factor	++	0	++		0	++	+	+
Thrombospondin 1	++	0	++		0	++	+	+
TIMP-2 (inhibitor of metalloproteinase 2)	++	+	++		+	++	+	+
TRIM-2/custerin (b)	+++	+	++		+	++	+	+



Sequence Identity (Genbank/EMBL)	Expression Strength			
	208F H-Ras	FE-8 K-Ras	208F N-Ras	208F N-Ras
ABC transporter MOAT-B	0	++++	0	+
BCSC-1 (breast cancer suppressor candidate 1)	+	++++	0	+
Cyclooxygenase 1	+	++++	+	+++
E1B 19K/Bcl-2-binding protein (Nip3)	0	++	++++	++
EST AA743557	++++	+	0	++
EST AA792426	+	++++	+	+
EST AA924000	+	++++	+	++
ETF TEA domain containing transcription factor	++++	0	++	++
Famesyl pyrophosphate synthetase	+	+++	0	+
FEN-1 (flap endonuclease-1)	0	++++	+	0
FLIP (FLICE-like inhibitory protein)	0	+	++	++++
JAK1 protein tyrosine kinase 1	+	++++	+	+
MAGE-B gene cluster	0	++++	0	0
MAP-kinase phosphatase (cpg21)	0	++	+++	++++
MARCKS	++++	0	+	+++
MMP-10 (Stromelysin 2)	0	++	++	++++
Mob-1 (<i>f</i>)	0	++++	++	+
mTFE3 (X-linked transcriptional activator)	++++	0	+	+
Myb-binding protein (P160)	+	++++	++	++
novel transcript N317	++++	0	++	++++
P-cadherin (<i>g</i>)	++++	0	0	++
Phosphatidylinositol 3-kinase p170	+++	0	+	++
Ras-GTPase-activating protein	0	++++	0	0
SBF1 phosphatase	0	++++	+	+
Serum inducible kinase (SNK) (<i>h</i>)	++++	0	+++	+++
Tyrosine phosphatase IA-2a (<i>i</i>)	0	++++	0	++

FIG. 4

**FIG. 5**

**FIG. 6**

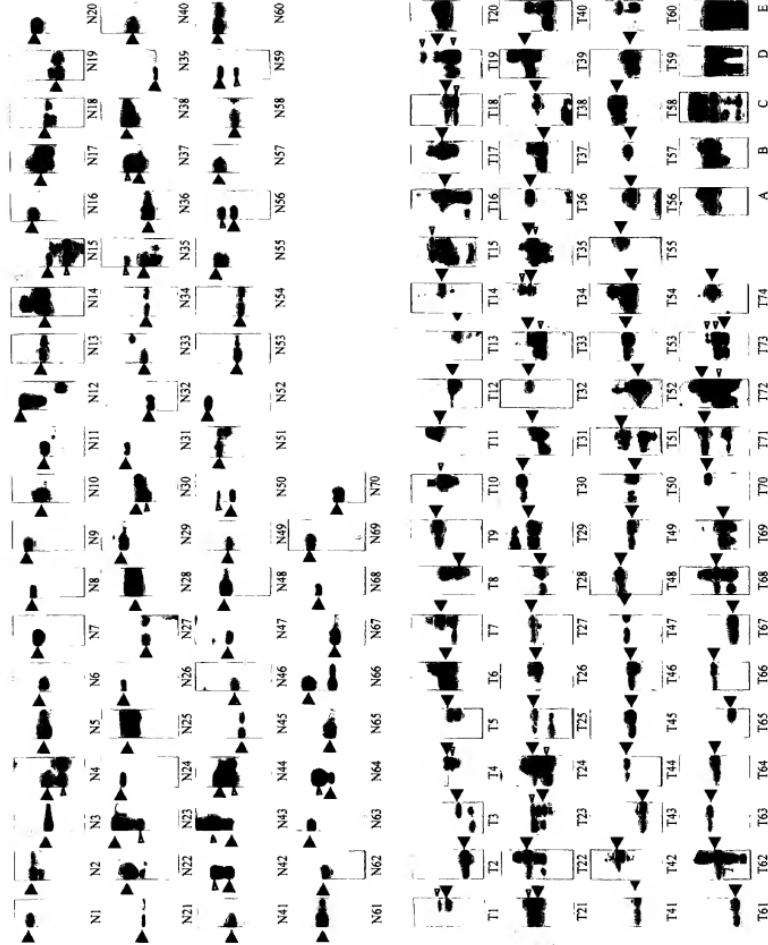


FIG. 7

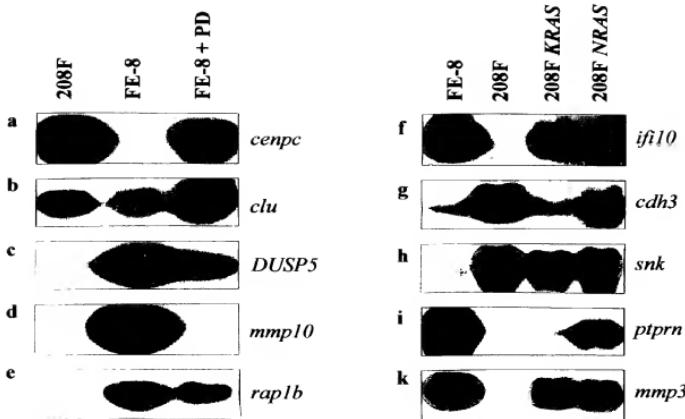


FIG. 8

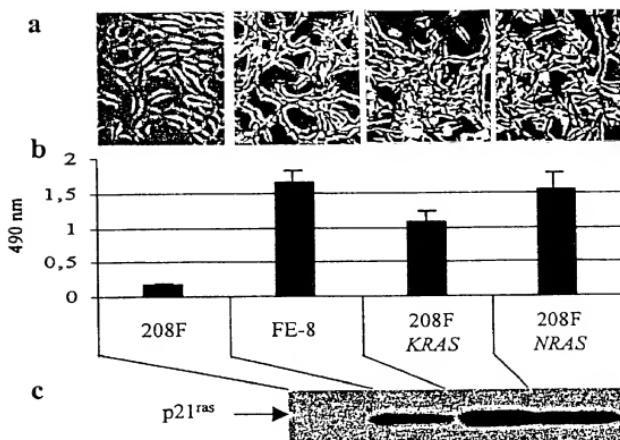


FIG. 9

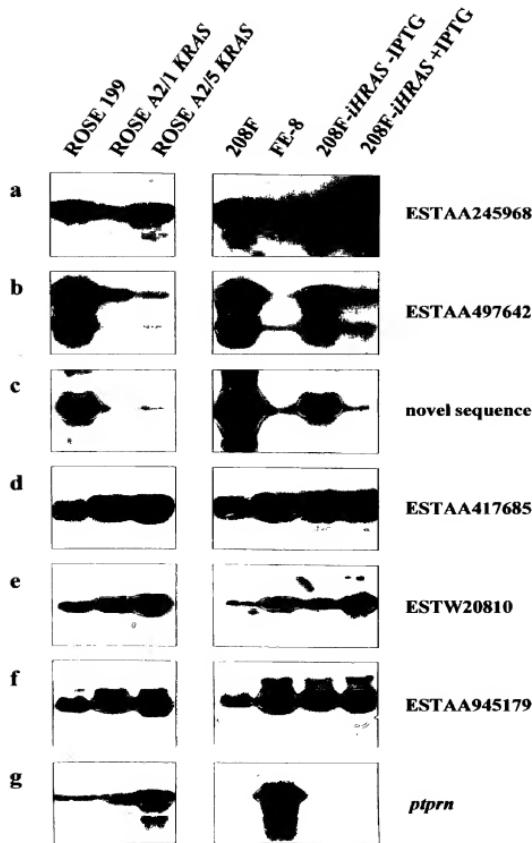


FIG. 10

1 T59
2 T182
3 T82
4 T6
5 T34
6 N5
7 N20
8 N280
9 N271
10 N126
11 T148
12 N199
13 T64
14 N131
15 T20
16 T162
17 T141
18 N77
19 N104
20 T49
21 T16
22 N189
23 N28
24 T124
25 T216
26 T60
27 T37
28 T160
29 N101
30 N40
31 T54
32 T120
33 N159
34 T185
35 N151
36 T147
37 N188
38 T25
39 T47
40 T43
41 T139
42 T176
43 N144
44 T35
45 T98
46 T15
47 T138
48 N21
49 T76
50 T103
51 T143
52 T44
53 N31
54 T243
55 N129
56 T193
57 T132
58 T137
59 T217
60 T191
61 N42
62 T156
63 T67

FIG. 11

64 N196
65 T21
66 N34
67 N134
68 T119
69 N36
70 N209
71 N256
72 T105
73 T75
74 T153
75 T189
76 T86
77 T111
78 T144
79 N192
80 N103
81 N270
82 N255
83 N61
84 N137
85 T174
86 N22
87 T2
88 T237
89 T19
90 N156
91 N59
92 N235
92 N248
92 N249
92 N252
92 N257
93 T38
94 T121
95 N10
96 T129
97 T66
98 T36
99 T40
100 N1
101 N212
102 T100
103 N112
104 N3
105 N238
106 T183
107 T238
108 T166
109 N29
110 T225
111 N175
112 N142
113 T72
114 N186
115 T212
116 T196
117 T48
118 N132
119 N158
120 T69
121 N7
122 T245

FIG. 11A

123 N102
124 T208
125 N44
126 T205
127 T215
128 N293
129 T226
130 T253
131 T222
132 N264
133 T240
134 N70
135 T125
136 N253
137 N234
138 N55
139 N202
140 N82
141 T45
142 T118
143 T10
144 N71
145 N183
146 N165
147 N213
148 N35
149 N182
150 N43
151 N75
152 T163
153 T89
154 N11
155 N32
156 T50
157 N215
158 N242
159 N181
160 N48
161 T227
162 N149
163 N109
164 N260
165 T219
166 T61
167 N85
168 N45
169 T250
170 N261
171 T172
172 N62
173 N160
174 N154
175 N58
176 T232
177 N128
178 N79
179 T58
180 N30
181 T68
182 T244
182 T251
182 T96
183 N26



FIG. 11B

184 N14
185 N121
186 T17
187 T3
188 T117
189 T14
190 T73
191 N4
192 N269
193 T239
194 T170
195 T146
196 N17
197 T235
198 N74
199 N18
200 T211
201 T186
201 T204
202 N50
203 N116
204 T223
205 N198
206 N267
207 T133
208 T80
209 N218
210 N266
211 T224
212 N148
213 N108
214 N263
215 N250
216 N92
217 N152
218 T11
219 T159
220 N243
221 N76
222 T116
223 T27
224 N207
225 T31
226 N38
227 N163
228 N81
229 T94
230 N228
231 N80
232 T230
233 T188
234 N180
235 N187
236 N136
237 N294
238 N275
239 N65
240 N89
241 N125
242 N205
243 N39
244 N13
245 T48



FIG. 11C

246 T100
247 T223
248 N104
249 N35
250 T245
251 N32
252 T62
253 N125
254 N180
255 N22
256 T61
257 T125
258 T174
259 T36
260 T19
261 T204
262 T153
263 T27
264 T212
265 T159
266 T226
267 T239
268 N263
269 T66
270 N75
271 N250
272 T175
273 N283
274 T40
275 N152
276 N256
277 N28
278 T160
279 T82
280 N122
281 T170
282 N44
283 N18
284 T103
285 N126
286 N55
287 T42
288 T34
289 N158
290 N21
291 N154
292 N80
293 T189
294 T17
295 T68
296 T14
297 T146
298 T120
299 N181
300 N192
301 T109
302 N215
303 T244
303 T251
304 T96
305 T211
306 T243
307 N218

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FIG. 11D



308	T224
309	T94
310	T183
311	N294
312	T191
313	T88
314	T9
315	N204
316	N175
317	N129
318	T141
319	N188
320	N209
321	T111
322	T144
323	N213
324	N109
325	N62
326	T235
327	N198
328	N148
329	N78
330	T116
331	N46
332	N49
333	N51
334	N52
335	T26

FIG. 11E